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RAW SEQUENCE LISTING

DATE: 08/07/2001

PATENT APPLICATION: US/09/911,569

TIME: 10:19:27

Input Set : A:\ES.txt

Output Set: N:\CRF3\08072001\1911569.raw

			SEQUENCE LISTING	
	4	(1) GENE	RAL INFORMATION:	
	6	(i)	APPLICANT: HAWLEY-NELSON, PAMELA	
	7		LAN, JIANQING	
	8		SHIH, POJEN	
	9		JESSE, JOEL A.	
,	10		SCHIFFERLI, KEVIN P.	
	11		GEBEYEHU, GULILAT	
•	13		TITLE OF INVENTION: PEPTIDE-ENHANCED TRANSFECTIONS	
	15	•	NUMBER OF SEQUENCES: 120	
	17	(iv)	CORRESPONDENCE ADDRESS:	
	18		(A) ADDRESSEE: GREENLEE, WINNER & SULLIVAN	
	19		(B) STREET: 5370 MANHATTAN CIRCLE, SUITE 201	
	20		(C) CITY: BOULDER	
	21		(D) STATE: CO	
	22	•	(E) COUNTRY: US	
	23	(\)	(F) ZIP: 80303	
	25	(∨)	COMPUTER READABLE FORM:	
	26 27		(A) MEDIUM TYPE: Floppy disk (B) COMPUTER: IBM PC compatible	
	28		(C) OPERATING SYSTEM: PC-DOS/MS-DOS	
	29		(D) SOFTWARE: PatentIn Release #1.0, Version #1.3	n
	31	(vi)	CURRENT APPLICATION DATA:	U
C>		(\ +)	(A) APPLICATION NUMBER: US/09/911,569	
C>			(B) FILING DATE: 23-Jul-2001	
• •	39		(C) CLASSIFICATION:	
C>		(vii)	PRIOR APPLICATION DATA:	
	37	(/	(A) APPLICATION NUMBER: US 09/039,780	
	38		(B) FILING DATE: 16-MAR-1998	
	41	(viii)	ATTORNEY/AGENT INFORMATION:	
	42		(A) NAME: SULLIVAN, SALLY A.	
	43		(B) REGISTRATION NUMBER: 32,064	٠.
	44		(C) REFERENCE/DOCKET NUMBER: 32-95D	
	46	(ix)	TELECOMMUNICATION INFORMATION:	
	48		(A) TELEPHONE: (303)499-8080	
. *	49		(B) TELEFAX: (303)499-8089	
			RMATION FOR SEQ ID NO: 1:	
	54	(i)	SEQUENCE CHARACTERISTICS:	
	55		(A) LENGTH: 10 amino acids	
	56		(B) TYPE: amino acid	
	57		(C) STRANDEDNESS: not relevant ()	
*	58		(D) TOPOLOGY: linear	
	60		MOLECULE TYPE: peptide	
	62		HYPOTHETICAL: NO	
	64		ANTI-SENSE: NO	
	69 71		SEQUENCE DESCRIPTION: SEQ ID NO: 1:	
	71	HIS	Ala Ile Arg Gly Asp Thr Phe Ala Thr	

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```
10
72
     . 1
74 (2) INFORMATION FOR SEQ ID NO: 2:
    (i) SEQUENCE CHARACTERISTICS:
            (A) LENGTH: 7 amino acids
77
78
            (B) TYPE: amino acid
        ___(C)_STRANDEDNESS: not relevant \bigcirc \checkmark
80
            (D) TOPOLOGY: linear
      (ii) MOLECULE TYPE: peptide
82
84
     (iii) HYPOTHETICAL: NO
      (iv) ANTI-SENSE: NO
86
      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
       Pro Lys Lys Lys Arg Lys Val
96 (2) INFORMATION FOR SEQ ID NO: 3:
       (i) SEQUENCE CHARACTERISTICS:
            (A) LENGTH: 8 amino acids
99
             (B) TYPE: amino acid
100
             (C) STRANDEDNESS: not relevant >
101
             (D) TOPOLOGY: linear
102
       (ii) MOLECULE TYPE: peptide.
104
       (iii) HYPOTHETICAL: NO
106
108
        (iv) ANTI-SENSE: NO
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
113
        Pro Lys Lys Lys Arg Lys Val
115
116
118 (2) INFORMATION FOR SEQ ID NO: 4:
120
        (i) SEQUENCE CHARACTERISTICS:
121
             (A) LENGTH: 20 amino acids
122
             (B) TYPE: amino acid
123
             (C) STRANDEDNESS: not relevant
124
             (D) TOPOLOGY: linear
126
       (ii) MOLECULE TYPE: peptide
128
       (iii) HYPOTHETICAL: NO
      (iv) ANTI-SENSE: NO
130
133
       (ix) FEATURE:
             (A) NAME/KEY: Modified-site 3
134
135
             (B) LOCATION: 2..20
             (D) OTHER INFORMATION: /product= "OTHER"
136
137 /note= "ANY OR ALL OF THE AMINO ACIDS 2-20 CAN BE PRESENT OR O
138 ABSENT"
141
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
143
        144.
                                                               15
        Lys Lys Lys
                    20
149 (2) INFORMATION FOR SEQ ID NO: 5:
151
        (i) SEQUENCE CHARACTERISTICS:
152
             (A) LENGTH: 20 amino acids
153
             (B) TYPE: amino acid
```

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```
PATENT APPLICATION: US/09/911,569
                Input Set : A:\ES.txt
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              (C) STRANDEDNESS: not relevant
154
              (D) TOPOLOGY: linear
155
        (ii) MOLECULE TYPE: peptide
157
160
        (ix) FEATURE:
161
              (A) NAME/KEY: Modified-site
162
              (B) LOCATION: 2...20
163
              (D) OTHER INFORMATION: /product= "OTHER"
164 /note= "ANY OR ALL OF THE AMINO ACIDS 2-20 CAN BE PRESENT OR
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
168
         170
171
                                             10
173
         Arg Arg Arg Arg
174
                     20
176 (2) INFORMATION FOR SEQ ID NO: 6:
        (i) SEQUENCE CHARACTERISTICS:
179
              (A) LENGTH: 40 amino acids
180
              (B) TYPE: amino acid
181
              (C) STRANDEDNESS: not relevant
182
              (D) TOPOLOGY: linear
184
        (ii) MOLECULE TYPE: peptide
187
        (ix) FEATURE:
              (A) NAME/KEY: Modified-site
188
189
              (B) LOCATION: 3..40
190
              (D) OTHER INFORMATION: /product= "OTHER"
191 /note= "ANY Lsy-Arg PAIR FROM AMINO ACID 3-40 CAN BE PRESENT OR
192 ABSENT"
195
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
197
         Lys Arg Lys Arg
198
                                             10
200
         Lys Arg Lys Arg Lys Arg Lys Arg Lys Arg Lys Arg Lys Arg
201
203
         Lys Arg Lys Arg Lys Arg Lys Arg
204
                 35
206 (2) INFORMATION FOR SEQ ID NO: 7:
208
         (i) SEQUENCE CHARACTERISTICS:
209
              (A) LENGTH: 20 amino acids
210
              (B) TYPE: amino acid
211
              (C) STRANDEDNESS: not relevant
212
              (D) TOPOLOGY: linear
214
        (ii) MOLECULE TYPE: peptide
217
        (ix) FEATURE:
218.
             (A) NAME/KEY: Modified-site
219
              (B) LOCATION: 2..20
              (D) OTHER INFORMATION: /product= "OTHER"
221 /note= "ANY OR ALL OF THE AMINO ACIDS 2-20 CAN BE PRESENT OR
222 ABSENT"
225
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
```

RAW SEQUENCE LISTING

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RAW SEQUENCE LISTING

DATE: 08/07/2001

PATENT APPLICATION: US/09/911,569 TIME: 10:19:27

Input Set : A:\ES.txt

Output Set: N:\CRF3\08072001\I911569.raw

```
15
                                          10
228
230
        Gly Gly Gly Gly
231
                    20
233 (2) INFORMATION FOR SEQ ID NO: 8:
235
        (i) SEQUENCE CHARACTERISTICS:
236
        (A) LENGTH: 7_amino_acids
237
             (B) TYPE: amino acid
238
             (C) STRANDEDNESS: not relevant
             (D) TOPOLOGY: linear
239
       (ii) MOLECULE TYPE: peptide
241
      (iii) HYPOTHETICAL: NO
243
       (iv) ANTI-SENSE: NO
245
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
250
        Gly Arg Gly Asp Ser Pro Cys
252
253
255 (2) INFORMATION FOR SEQ ID NO: 9:
257
        (i) SEQUENCE CHARACTERISTICS:
258
             (A) LENGTH: 23 amino acids
259
             (B) TYPE: amino acid
260
             (C) STRANDEDNESS: not relevant
261
             (D) TOPOLOGY: linear
       (ii) MOLECULE TYPE: peptide
263
       (ix) FEATURE:
266
             (A) NAME/KEY: Modified-site
267
268
             (B) LOCATION: 1..20
             (D) OTHER INFORMATION: /product= "OTHER"
269
270 /note= "ANY OR ALL OF THE AMINO ACIDS 1-20 CAN BE PRESENT OR
271 ABSENT"
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
276
        277
        1
279
        Lys Lys Lys Arg Gly Asp
280
                    20
282 (2) INFORMATION FOR SEQ ID NO: 10:
        (i) SEQUENCE CHARACTERISTICS:
284
285
             (A) LENGTH: 19 amino acids
286
             (B) TYPE: amino acid
287
             (C) STRANDEDNESS: not relevant
288
             (D) TOPÖLOGY: linear
290
       (ii) MOLECULE TYPE: peptide
295
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
297
        298
                                          10
                                                             15
300
        Arg Gly Asp
303 (2) INFORMATION FOR SEQ ID NO: 11:
305
        (i) SEQUENCE CHARACTERISTICS:
306
             (A) LENGTH: 7 amino acids
307
             (B) TYPE: amino acid
308
             (C) STRANDEDNESS: not relevant
```

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RAW SEQUENCE LISTING
                                                          DATE: 08/07/2001
                                                          TIME: 10:19:27
                PATENT APPLICATION: US/09/911,569
                Input Set : A:\ES.txt
                Output Set: N:\CRF3\08072001\1911569.raw
309
               (D) TOPOLOGY: linear
        (ii) MOLECULE TYPE: peptide
311
313
       (iii) HYPOTHETICAL: NO
        (iv) ANTI-SENSE: NO
315
320
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
         Lys Phe Thr Ile Val Phe Cys
322
323
                          5
325 (2) INFORMATION FOR SEQ ID NO: 12:
         (i) SEQUENCE CHARACTERISTICS:
327
328
               (A) LENGTH: 12 amino acids
329
              (B) TYPE: amino acid
330
              (C) STRANDEDNESS: not relevant
331
              (D) TOPOLOGY: linear
333
        (ii) MOLECULE TYPE: peptide
335
       (iii) HYPOTHETICAL: NO
337
        (iv) ANTI-SENSE: NO
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:
342
344
         Cys Gly Trp Gly Pro Lys Lys Lys Arg Lys Val Gly
345
         1
                          5
                                               10
347 (2) INFORMATION FOR SEQ ID NO: 13:
         (i) SEQUENCE CHARACTERISTICS:
350
              (A) LENGTH: 26 amino acids
351
              (B) TYPE: amino acid
352
              (C) STRANDEDNESS: not relevant
353
              (D) TOPOLOGY: linear
355
        (ii) MOLECULE TYPE: peptide
357
       (iii) HYPOTHETICAL: NO
359
        (iv) ANTI-SENSE: NO
364
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:
366
         Gly Gly Tyr Gly Pro Lys Lys Lys Arg Lys Val Gly Gly Gly Tyr
367
369
         Gly Pro Lys Lys Lys Arg Lys Val Gly Gly
370
                      20
372 (2) INFORMATION FOR SEQ ID NO: 14:
374
         (i) SEQUENCE CHARACTERISTICS:
375
              (A) LENGTH: 6 amino acids
376
              (B) TYPE: amino acid.
377
              (C) STRANDEDNESS: not relevant
              (D) TOPOLOGY: linear
378
380
        (ii) MOLECULE TYPE: peptide
382
       (iii) HYPOTHETICAL: NO
384
        (iv) ANTI-SENSE: NO
389
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:
391
         Arg Gly Asp Ser Pro Cys
392
         1
```

Please Note:

396

397

398 -

394 (2) INFORMATION FOR SEQ ID NO: 15:

(i) SEQUENCE CHARACTERISTICS:

(B) TYPE: amino acid

(A) LENGTH: 4 amino acids

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a c rresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/911,569

L:32 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]

DATE: 08/07/2001 TIME: 10:19:28

Input Set : A:\ES.txt

Output Set: N:\CRF3\08072001\I911569.raw

```
L:33 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
L:36 M:220 C: Keyword misspelled or invalid format, [(vii) PRIOR APPLICATION DATA:]
L:553 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
L:633 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L:676 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:679 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:682 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:746 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:749 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:752 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:755 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:758 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:761 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:817 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:820 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:823 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:826 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:829 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:892 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:895\ M:341\ W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:898 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:901 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:904 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:907 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:970 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 L:973 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:976 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:979 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:982 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:985 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:1048 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30
L:1051 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30
L:1054 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30
L:1057 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30
L:1060 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30
L:1063 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30
L:1143 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32
L:1146 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32
L:1149 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32
L:1152 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32
L:1155 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32
L:1216 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33
L:1219 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33
L:1222 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33
L:1225 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 L:1228 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 L:1290 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34
```

VERIFICATION SUMMARY

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Input Set : A:\ES.txt

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L:1293 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 L:1296 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 $L:1299 \ M:341 \ W: (46)$ "n" or "Xaa" used, for SEQ ID#:34 L:1302 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34
L:1364 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35
L:5082 M:246 W: Invalid value of Alpha Sequence Header Field, [TOPOLOGY:], SeqNo=115